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U.S. PATENT AND TRADEMARK OFFICE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/020,721

DATE: 01/07/2002
TIME: 20:35:24

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\01072002\J020721.raw

3 <110> APPLICANT: HITACHI SOFTWARE ENGINEERING CO., LTD.
5 <120> TITLE OF INVENTION: DETECTION METHOD AND DETECTION KIT FOR PCR AMPLIFIED
6 BASE SEQUENCES
8 <130> FILE REFERENCE: PH-1431
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/020,721
C--> 11 <141> CURRENT FILING DATE: 2001-12-14
13 <160> NUMBER OF SEQ ID NOS: 7
15 <170> SOFTWARE: PatentIn Ver. 2.0
17 <210> SEQ ID NO: 1
18 <211> LENGTH: 336
19 <212> TYPE: DNA
20 <213> ORGANISM: Hepatitis C virus
22 <400> SEQUENCE: 1
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24 accttctcgc ctcgcggca tgagacagt caggactgca actgctcaat ctaccccgcc 120
25 catgtatcag gtcacccat ggcttggat atgatgatga actggtcacc tacaacagcc 180
26 ctatgggtgt cgcagggtct ccggatccca caagctgtcg tggacatgtt ggcggggggcc 240
27 cactggggag tcctggcggg cttgcctac tattccatgg tagggactg ggctaagggtt 300
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33 <211> LENGTH: 336
34 <212> TYPE: DNA
35 <213> ORGANISM: Hepatitis C virus
37 <400> SEQUENCE: 2
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39 accttctcac ctcgcggta tgagacggta caagactgca attgctcaat ctatcccgcc 120
40 cacgtatcag gtcacccat ggcttggat atgatgatga actggtcacc tacaacagcc 180
41 ctgggtgtat cgcagggtct ccggatccca caagccgtcg tggacatgtt ggcggggggcc 240
42 cactggggag tcctagcggg cttgcctac tattccatgg tggggactg ggctaagggtt 300
43 ctgattgtga tgctactt tgccggcggtt gacggg 336
45 <210> SEQ ID NO: 3
46 <211> LENGTH: 336
47 <212> TYPE: DNA
48 <213> ORGANISM: Hepatitis C virus
50 <400> SEQUENCE: 3
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53 catataacgg gtcatcgat gcatggat atgatgatga actggccccc tacggcagcg 180
54 ttgggtgtat ctcagctgtt ccggatccca caagccatca tggacatgtt cgctgggtct 240
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56 ctggtagtgc tgctgtatt tgccggcggtt gacggt 336
58 <210> SEQ ID NO: 4
59 <211> LENGTH: 19
60 <212> TYPE: DNA
61 <213> ORGANISM: Hepatitis C virus
65 <400> SEQUENCE: 4
66 tgctccggat cccacaaggc

19

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Input Set : A:\PTO.AMC.txt
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68 <210> SEQ ID NO: 5
69 <211> LENGTH: 20
70 <212> TYPE: DNA
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77 <211> LENGTH: 20
78 <212> TYPE: DNA
79 <213> ORGANISM: Hepatitis C virus
81 <400> SEQUENCE: 6 20
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84 <210> SEQ ID NO: 7
85 <211> LENGTH: 20
86 <212> TYPE: DNA
87 <213> ORGANISM: Hepatitis C virus
89 <400> SEQUENCE: 7 20
90 ttcttgg tcaactgttt

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/020,721

DATE: 01/07/2002

TIME: 20:35:25

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\01072002\J020721.raw

L:10 M:270 C: Current Application Number differs, Replaced Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/020,721

DATE: 01/02/2002
TIME: 10:26:07

Input Set : A:\pto.vsk.txt
Output Set: N:\CRF3\01022002\J020721.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: HITACHI SOFTWARE ENGINEERING CO., LTD.
5 <120> TITLE OF INVENTION: DETECTION METHOD AND DETECTION KIT FOR PCR AMPLIFIED
6 BASE SEQUENCES
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C--> 11 <141> CURRENT FILING DATE: 2001-12-14
13 <160> NUMBER OF SEQ ID. NOS: 7
15 <170> SOFTWARE: PatentIn Ver. 2.0

ERRORED SEQUENCES

84 <210> SEQ ID NO: 7
85 <211> LENGTH: 20
86 <212> TYPE: DNA
87 <213> ORGANISM: Hepatitis C virus
89 <400> SEQUENCE: 7
90 ttcttggttgg tcaactgttt
E--> 92 (1/1)

20

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/020,721

DATE: 01/02/2002
TIME: 10:26:08

Input Set : A:\pto.vsk.txt
Output Set: N:\CRF3\01022002\J020721.raw

L:10 M:270 C: Current Application Number differs, Replaced Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:92 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:21 SEQ:7
L:92 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:92 M:252 E: No. of Seq. differs, <211>LENGTH:Input:20 Found:21 SEQ:7

1645 050 000 01RE

CRF Errors Corrected by the STIC Sy

Serial Number: 10/020,721

ns Branch

CRF Processing Date:

Edited by:

Verified by:

10/7/2002
2 8 2002
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ENTERED

Changed a file from non-ASCII to ASCII

Changed the margins in cases where the sequence text was 'wrapped' down to the next line.

Edited a format error in the Current Application Data section, specifically:

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other _____

Added the mandatory heading and subheadings for 'Current Application Data'.

Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integer.

Changed the spelling of a mandatory field (the headings or subheadings), specifically:

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included: _____

Deleted extra, invalid, headings used by an applicant, specifically:

Deleted: non-ASCII 'garbage' at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as _____

Inserted mandatory headings, specifically: _____

Corrected an obvious error in the response, specifically:

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically:

A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be deleted.

Deleted ending stop codon in amino acid sequences and adjusted the '(A)Length:' field accordingly (error due to a PatentIn bug). Sequences corrected: _____

Other:

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1995